

DEX-0275

AA

ID AQ743305 standard; DNA; GSS; 1141 BP.

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AC AQ743305;

XX XP-002214818

SV AQ743305.1

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DT 19-JUL-1999 (Rel. 60, Created)

DT 04-MAR-2000 (Rel. 63, Last updated, Version 2)

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DE HS\_5387\_B2\_B02\_SP6 RPCI-11 Human Male BAC Library Homo sapiens genomic

DE clone Plate=963 Col=4 Row=D, genomic survey sequence.

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KW GSS.

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OS Homo sapiens (human)

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;

OC Eutheria; Primates; Catarrhini; Hominidae; Homo.

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RN [1]

RP 1-1141

RX MEDLINE; 99380589.

RA Mahairas G.G., Wallace J.C., Smith K., Swartzell S., Holzman T., Keller A.,

RA Shaker R., Furlong J., Young J., Zhao S., Adams M.D., Hood L.;

RT "Sequence-tagged connectors: A sequence approach to mapping and scanning  
the human genome";

RL Proc. Natl. Acad. Sci. U.S.A. 96(17):9739-9744 (1999).

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CC High Throughput Sequencing Center

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CC Clones are derived from the human BAC library RPCI-11. For BAC

CC library availability, please contact Pieter de Jong

CC (pieter@dejong.med.buffalo.edu). Clones may be purchased from

CC BACPAC Resources ([http://bacpac.med.buffalo.edu/ordering\\_bac.htm](http://bacpac.med.buffalo.edu/ordering_bac.htm))CC or from Research Genetics ([info@resgen.com](mailto:info@resgen.com)). BAC end Web Server:CC <http://www.htsc.washington.edu>

CC Plate: 963 row: D column: 4

CC Seq primer: SP6

CC Class: BAC ends

CC High quality sequence stop: 1141.

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FH Key Location/Qualifiers

FH

FT source 1..1141

FT /db\_xref="taxon:9606"

FT /note="Vector: pBACe3.6; Genomic sequence of BAC ends"

FT /sex="male"

FT /organism="Homo sapiens"

FT /clone="Plate=963 Col=4 Row=D"

FT /clone\_lib="RPCI-11 Human Male BAC Library"

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SQ Sequence 1141 BP; 760 A; 301 C; 16 G; 15 T; 49 other;

aacgttaaggc aactnnnaag annaagacna tnnanctnnn nnnagactnn nccctcntaa	60
attnnaaggg cggccaanaa anccctttan aaggaaaaan tacanggacc ncnnaaaaana	120
nannaaaaaaa aaaaaacaaa caaaaaattta aaaaacaaaaa aaaaaaaaaa ccacaaaaaaa	180
cacacacaac naaaacacaa aaaaacaaaaa aacaaaccac aacaacacaa caaaaaaccca	240
aaacaaaccaa aaaaacaaca aaaaaaaaaa aaaaaaaaaca aaaaaaaaaa ccanacacac	300
aacaacacaa caacaacacc aaaaacacaca cacaacacnc accaaaaaaaaa caacaacacaa	360
aaacacaaca aaaaacaaac aaaccacgac accaaacaac aacacacaca aaaaacacaac	420
aaaaaaaaaaa aaaaaccaac aaacaaccaa aaaaaaaaaaac caaaaccaac aacacacacaa	480

P.D. 19-07-1999	2
P. 1-2	

caacaccaac aacaacaaaa caacaaaaca	aaaacaacaa	caaaccacaa	aaaaaacaaa	540
cacacaacaa ncacaaaaaa	acaacaaaac	aaaacaacaa	aaaaacacaa	600
acaaaaacaa caaacaacac	aaccacacaa	aacaacaaca	acaacaacaa	660
ananaaacac acaaaaaaca	aacaaaaaaa	acaaccaaca	aaaaacacaa	720
aaaaaaaaacc aaaaaaacac	aancacacaa	cccacaaaac	acacaaaaaa	780
cacacaacac aacaaccaa	aaaaaaccac	acacacaaac	caacaaaaca	840
caaaccacac caacaaaacc	acacaaacaa	ancaacaaca	caaacaacaa	900
cacaccacca caacacaca	caacaanaaa	accacaaaaa	cacacaaaac	960
acaaaaaccaa ancaaccaac	aacaaaaaac	aacaacaac	acaacaaaaa	1020
acaaaaaana ccaaacaac	aaacaacaaa	caccacaaca	accaacaaaa	1080
aaacaaaaaa acaaaaaaaa	aaaaacacaa	aaaaacaaca	caacacaaaa	1140
			aacacaaanc	1141

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